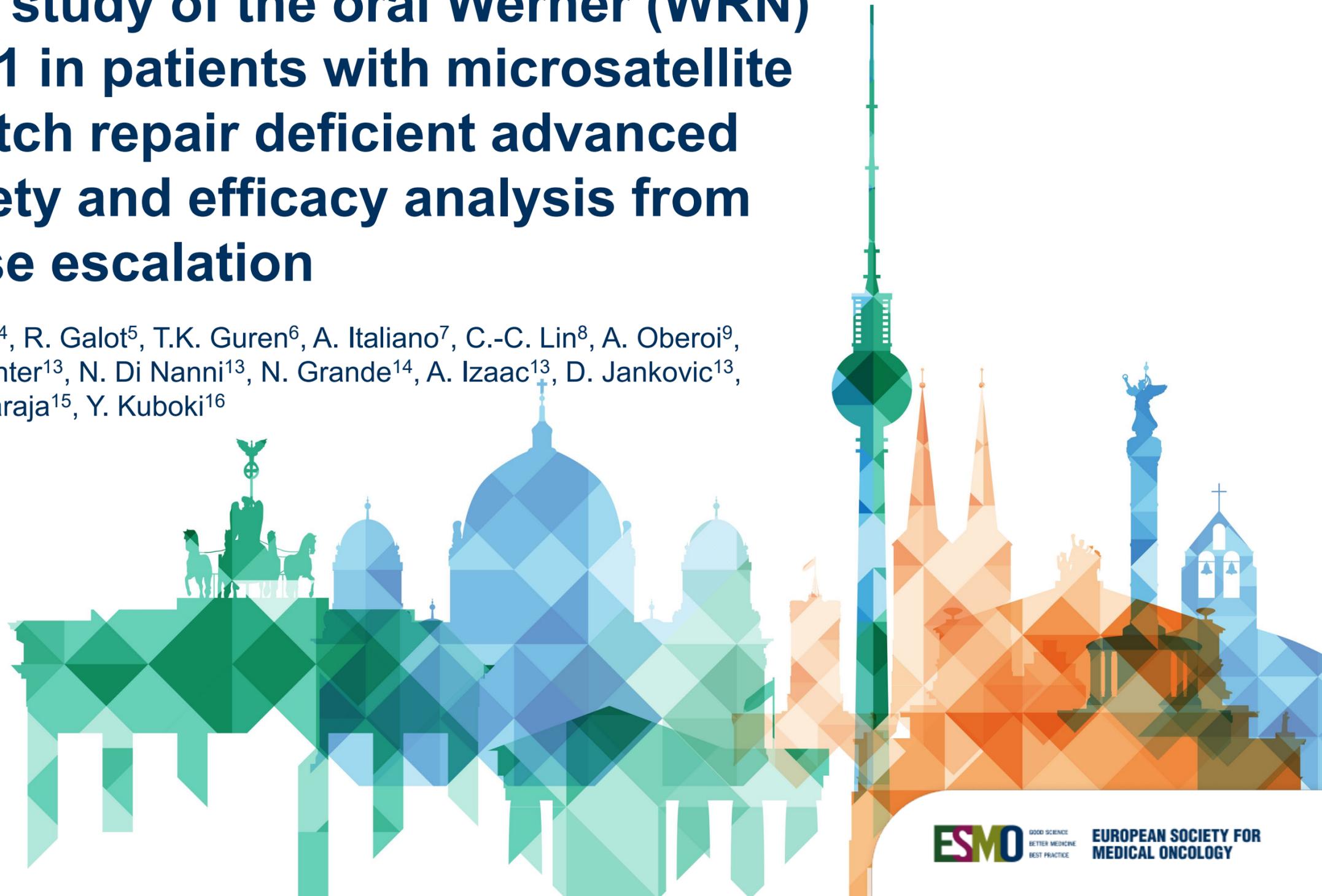


# First-in-human phase I/Ib study of the oral Werner (WRN) helicase inhibitor HRO761 in patients with microsatellite instability-high or mismatch repair deficient advanced solid tumors: Interim safety and efficacy analysis from HRO761 single agent dose escalation

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17 October 2025



# Declaration of interests

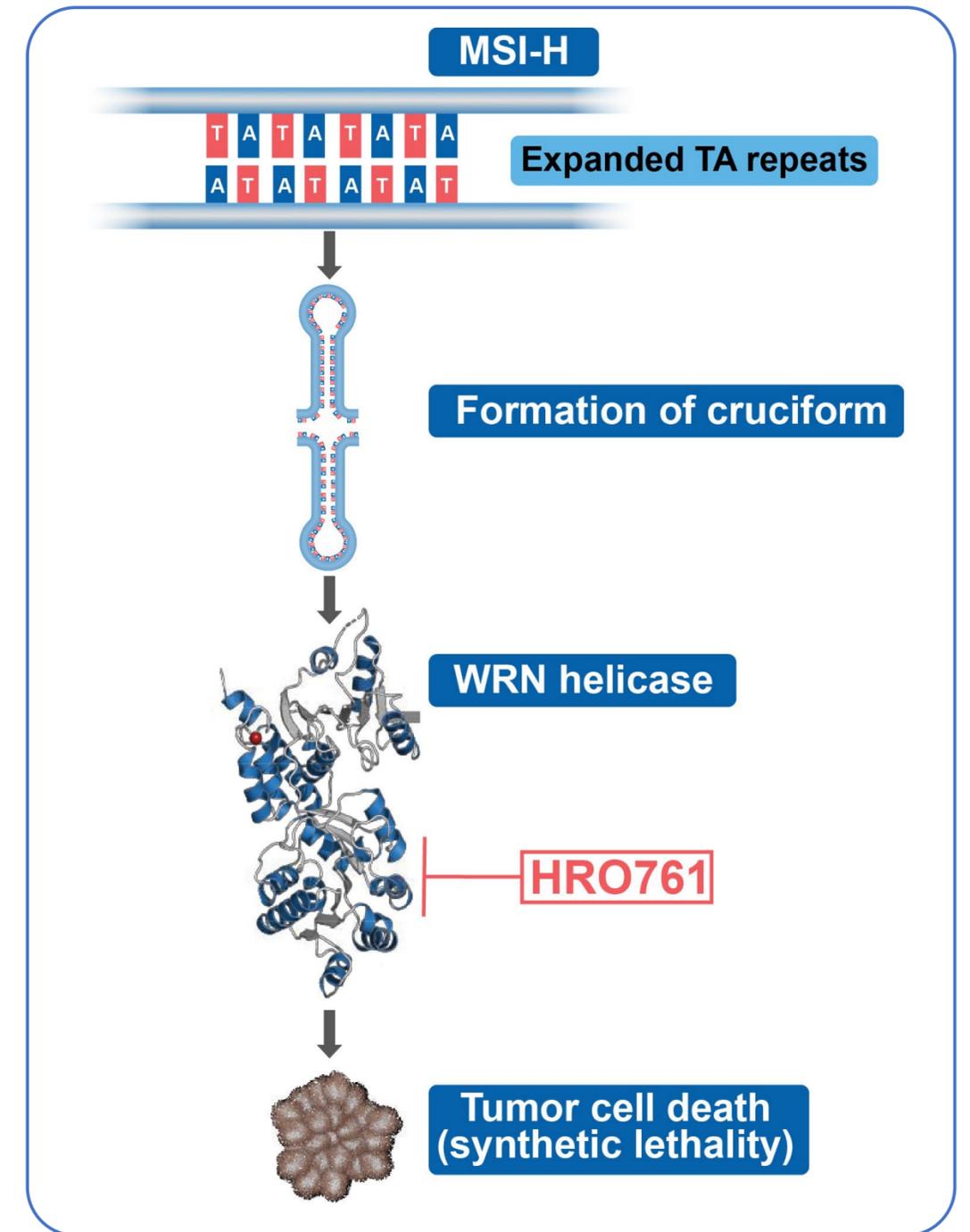
**Michael B. Foote**

**Advisory board, personal:** Bristol Meyers Squibb, Genzyme, Abbott Laboratories, Intera

**Medical Education Talk, personal:** Axiom Healthcare Strategies, Cancer Network, Horizon CME, OncLive

# HRO761 is an oral WRN inhibitor for MSI-H cancers

- MSI-H cancers exhibit secondary structures of expanded TA-dinucleotide repeats that require unwinding by the WRN syndrome protein, a DNA helicase<sup>1</sup>
- WRN inhibition in MSI-H cells induces synthetic lethality through DNA damage, cell cycle arrest, and apoptosis<sup>1,2</sup>
- HRO761 is a potent, selective, allosteric WRN inhibitor that locks WRN in an inactive conformation<sup>3</sup>
- HRO761 induces strong, dose-dependent regression of tumors in MSI-H cell- and patient-derived xenograft models<sup>3</sup>



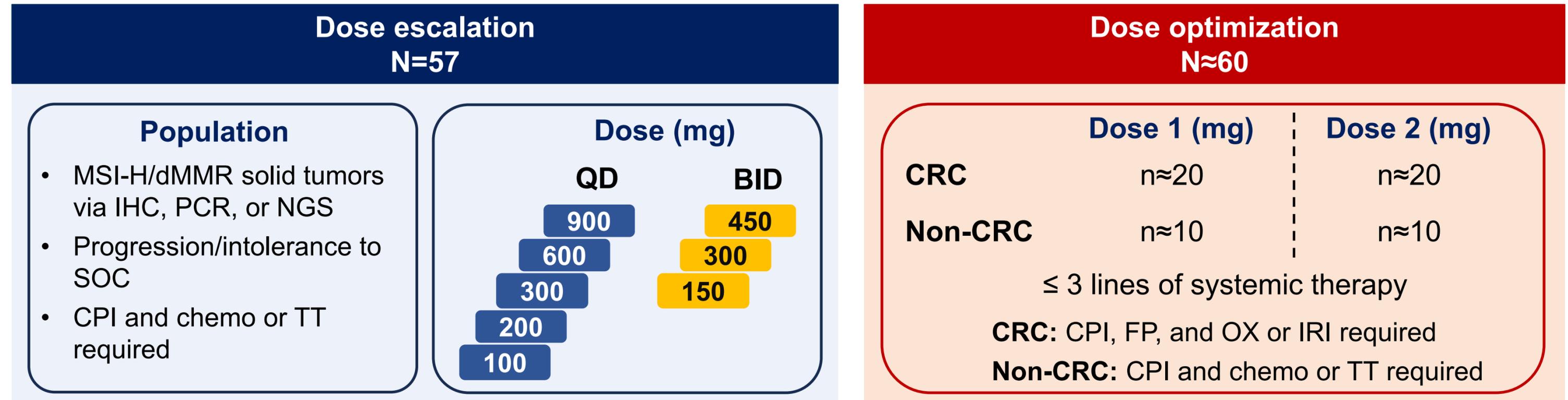
MSI-H, microsatellite instability-high; WRN, Werner.

1. van Wietmarschen N, et al. *Nature*. 2020;586:292–298; 2. Chan EM, et al. *Nature*. 2019;568:551–556; 3. Ferretti S, et al. *Nature*. 2024;629:443–449. 4. Moschetta M, et al. *ESMO Congress*. 2023; Poster 719TiP.

Adapted from Moschetta M, et al.<sup>4</sup>

# Phase I dose finding of single agent HRO761 in patients with MSI-H/dMMR advanced solid tumors

Data cutoff: June 15, 2025



- |   |  |
|---|--|
| <h3 style="color: #002060;">Primary objectives</h3> <ul style="list-style-type: none"> <li>Safety and tolerability</li> <li>Recommended dose and regimen</li> </ul> | <h3 style="color: #002060;">Secondary objectives</h3> <ul style="list-style-type: none"> <li>PK profile</li> <li>Preliminary antitumor activity per RECIST v1.1</li> </ul> |
|---|--|

EU-CT number: 2022-502314-93; ClinicalTrials.gov ID: NCT05838768  
 BID, twice daily; chemo, chemotherapy; CPI, checkpoint inhibitor; CRC, colorectal cancer; dMMR, mismatch repair deficient; EU-CT, European Union-Clinical Trial; FP, fluoropyrimidine; ID, identifier; IHC, immunohistochemistry; IRI, irinotecan; MSI-H, microsatellite instability-high; NGS, next-generation sequencing; OX, oxaliplatin; PCR, polymerase chain reaction; PK, pharmacokinetics; QD, once daily; RECIST v1.1, Response Evaluation Criteria in Solid Tumors version 1.1; SOC, standard of care; TT, targeted therapy.

# Baseline characteristics

Baseline characteristics	All patients (N=57)
Age (years), median (range)	56.0 (32–85)
Sex, n (%) Female / Male	29 (50.9) / 28 (49.1)
Race, n (%) Asian / Black or African American / White / NR or UNK	22 (38.6) / 2 (3.5) / 26 (45.6) / 7 (12.3)
ECOG PS, n (%) 0 / 1	18 (31.6) / 39 (68.4)
Prior antineoplastic therapy, n (%) Number of regimens, n (%) 1 / 2	57 (100) 2 (3.5) / 9 (15.8)
≥ 3 Median (range)	46 (80.7) 4 (1–10)
Initial diagnosis, n (%)	
CRC*	33 (57.9)
Non-CRC	24 (42.1)
Endometrial*	6 (10.5)
Gastric/GEJ*	6 (10.5)
Other	12 (21.1)

\*Adenocarcinoma.

CRC, colorectal cancer; ECOG PS, Eastern Cooperative Oncology Group performance status; GEJ, gastroesophageal junction; NR, not reported; UNK, unknown.

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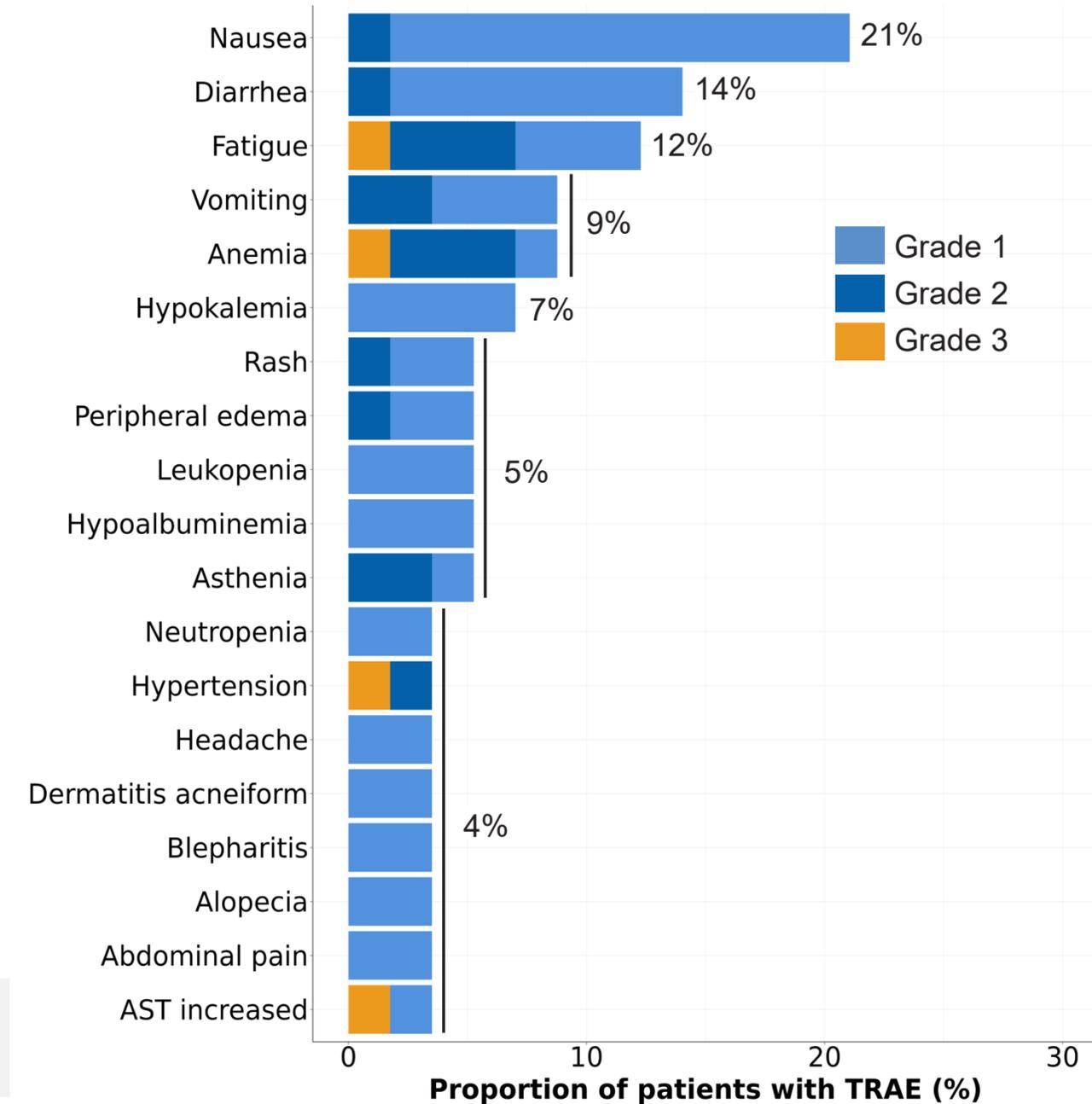
# HRO761 shows a favorable safety profile

## Overview of all AEs by dose and regimen

	100 mg QD (N=8)	200 mg QD (N=10)	300 mg QD (N=4)	150 mg BID (N=6)	600 mg QD (N=10)	300 mg BID (N=5)	900 mg QD (N=9)	450 mg BID (N=5)	All (N=57)
<b>AEs, n (%)</b>	7 (87.5)	9 (90.0)	4 (100)	5 (83.3)	10 (100)	5 (100)	8 (88.9)	5 (100)	<b>53 (93.0)</b>
<b>Treatment-related</b>	<b>3 (37.5)</b>	<b>6 (60.0)</b>	<b>3 (75.0)</b>	<b>4 (66.7)</b>	<b>7 (70.0)</b>	<b>4 (80.0)</b>	<b>5 (55.6)</b>	<b>2 (40.0)</b>	<b>34 (59.6)</b>
<b>AEs with grade ≥3, n (%)</b>	4 (50.0)	6 (60.0)	2 (50.0)	3 (50.0)	3 (30.0)	4 (80.0)	4 (44.4)	0	<b>26 (45.6)</b>
<b>Treatment-related</b>	<b>1 (12.5)</b>	<b>2 (20.0)</b>	<b>0</b>	<b>0</b>	<b>0</b>	<b>2 (40.0)</b>	<b>3 (33.3)</b>	<b>0</b>	<b>8 (14.0)</b>
<b>SAEs, n (%)</b>	3 (37.5)	5 (50.0)	2 (50.0)	3 (50.0)	3 (30.0)	2 (40.0)	2 (22.2)	0	<b>20 (35.1)</b>
<b>Treatment-related</b>	<b>0</b>	<b>1 (10.0)</b>	<b>0</b>	<b>0</b>	<b>0</b>	<b>1 (20.0)</b>	<b>1 (11.1)</b>	<b>0</b>	<b>3 (5.3)</b>
<b>AEs leading to dose interruption, n (%)</b>	4 (50.0)	5 (50.0)	1 (25.0)	3 (50.0)	4 (40.0)	4 (80.0)	5 (55.6)	2 (40.0)	<b>28 (49.1)</b>
<b>Treatment-related</b>	<b>2 (25.0)</b>	<b>2 (20.0)</b>	<b>0</b>	<b>2 (33.3)</b>	<b>1 (10.0)</b>	<b>2 (40.0)</b>	<b>2 (22.2)</b>	<b>0</b>	<b>11 (19.3)</b>
<b>AEs leading to dose adjustment, n (%)</b>	0	0	0	0	1 (10.0)	0	2 (22.2)	0	<b>3 (5.3)</b>
<b>Treatment-related</b>	<b>0</b>	<b>0</b>	<b>0</b>	<b>0</b>	<b>1 (10.0)</b>	<b>0</b>	<b>1 (11.1)</b>	<b>0</b>	<b>2 (3.5)</b>
<b>AEs leading to treatment discontinuation, n (%)</b>	1 (12.5)	1 (10.0)	1 (25.0)	0	2 (20.0)	0	0	0	<b>5 (8.8)</b>
<b>Treatment-related</b>	<b>0</b>	<b>0</b>	<b>0</b>	<b>0</b>	<b>0</b>	<b>0</b>	<b>0</b>	<b>0</b>	<b>0</b>
<b>DLT</b>	0	0	0	0	0	0	1 (11.1)	0	<b>1 (1.8)</b>

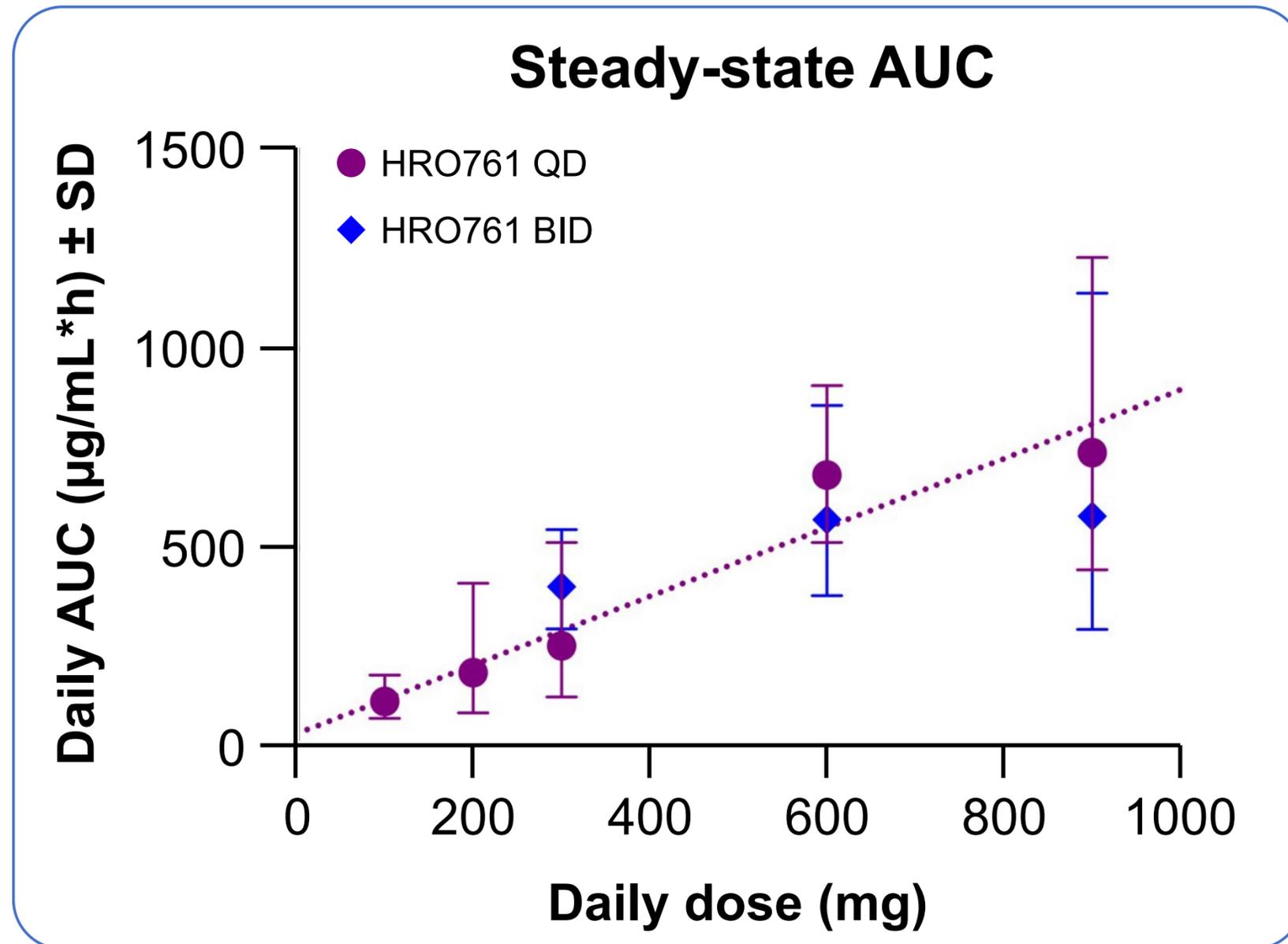
- Most common TRAEs were low-grade nausea, diarrhea, and fatigue
- One DLT of decreased EF without recurrence after dose reduction from 900 to 600 mg QD

## TRAEs occurring in 2 or more patients (N=57)



AE, adverse event; AST, aspartate aminotransferase; BID, twice daily; DLT, dose-limiting toxicity; EF, ejection fraction; QD, once daily; SAE, serious adverse event; TRAE, treatment-related adverse event.

# HRO761 shows a dose-linear increase in exposure without differences in QD and BID schedules

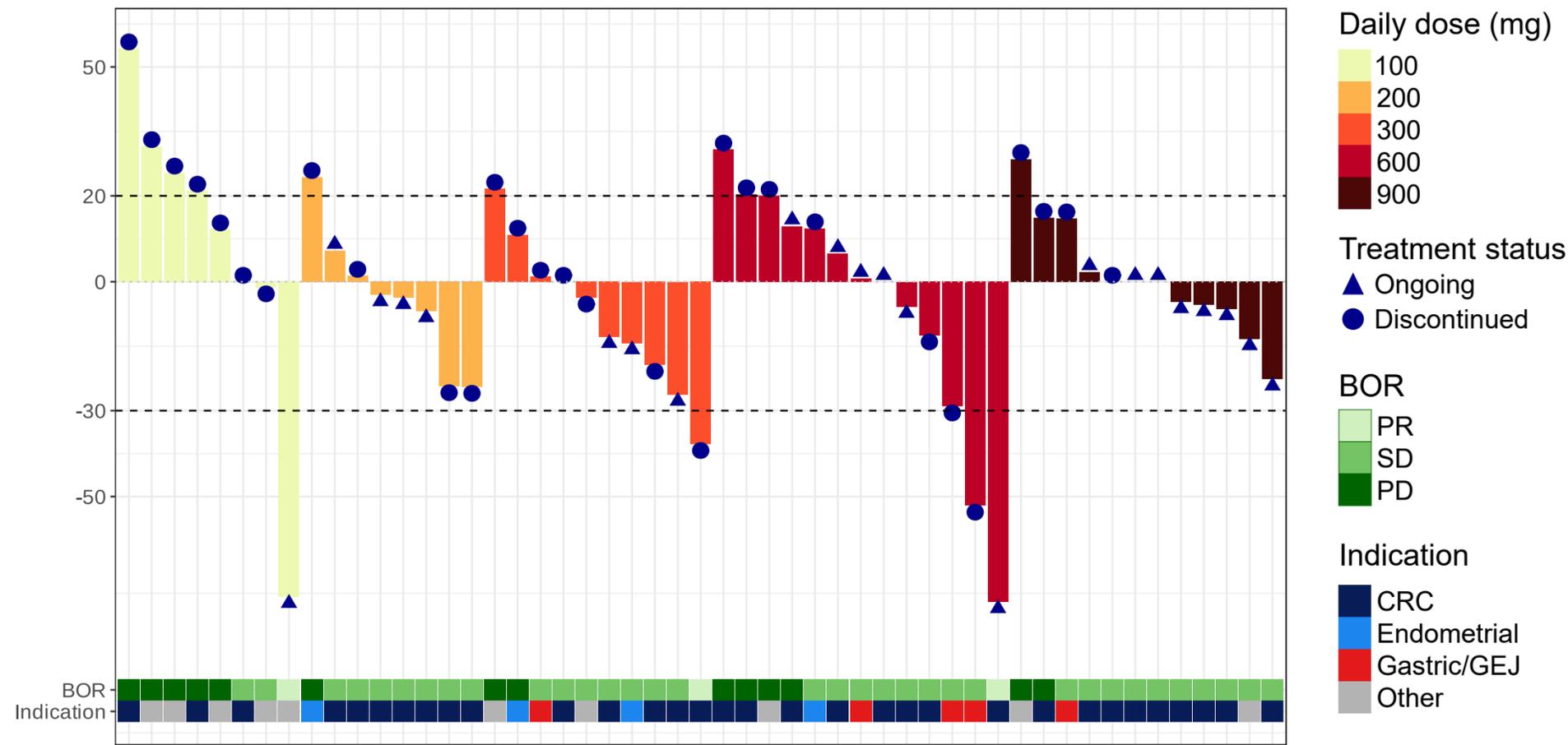


- QD dosing shows similar exposure as BID dosing, at equal total daily dose
- 2.1-fold accumulation was observed after repeated dosing on day 8 vs. day 1 in cycle 1
- Effective half-life = 19 hours
- Dose-linear increase in exposure up to 900 mg/day

Geometric mean plasma AUC (cycle 1, day 8) is shown.  
AUC, area under the curve; BID, twice daily; QD, once daily; SD, standard deviation.

# HRO761 shows antitumor activity with a high disease control rate

## Best percent change in tumor lesions



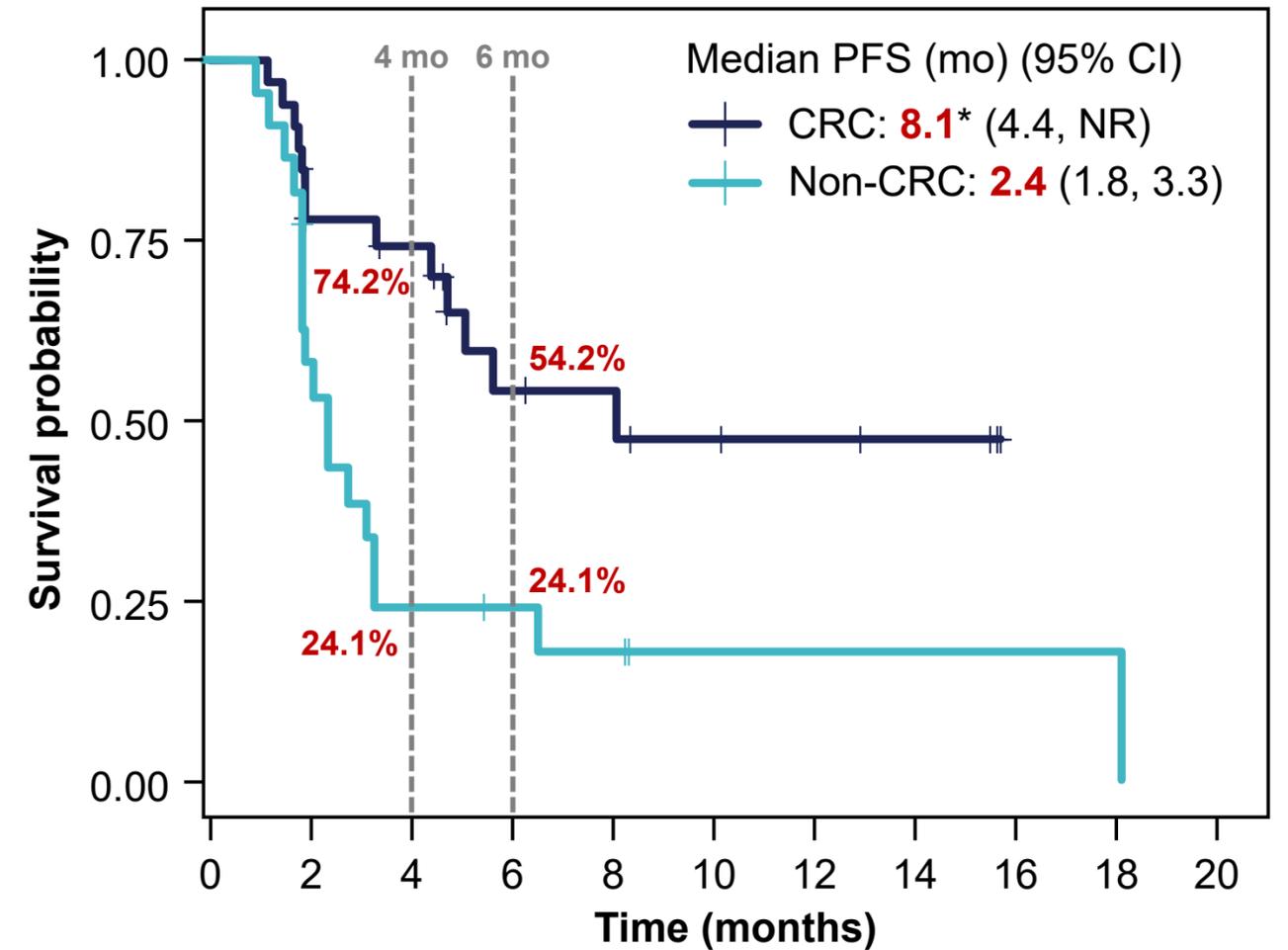
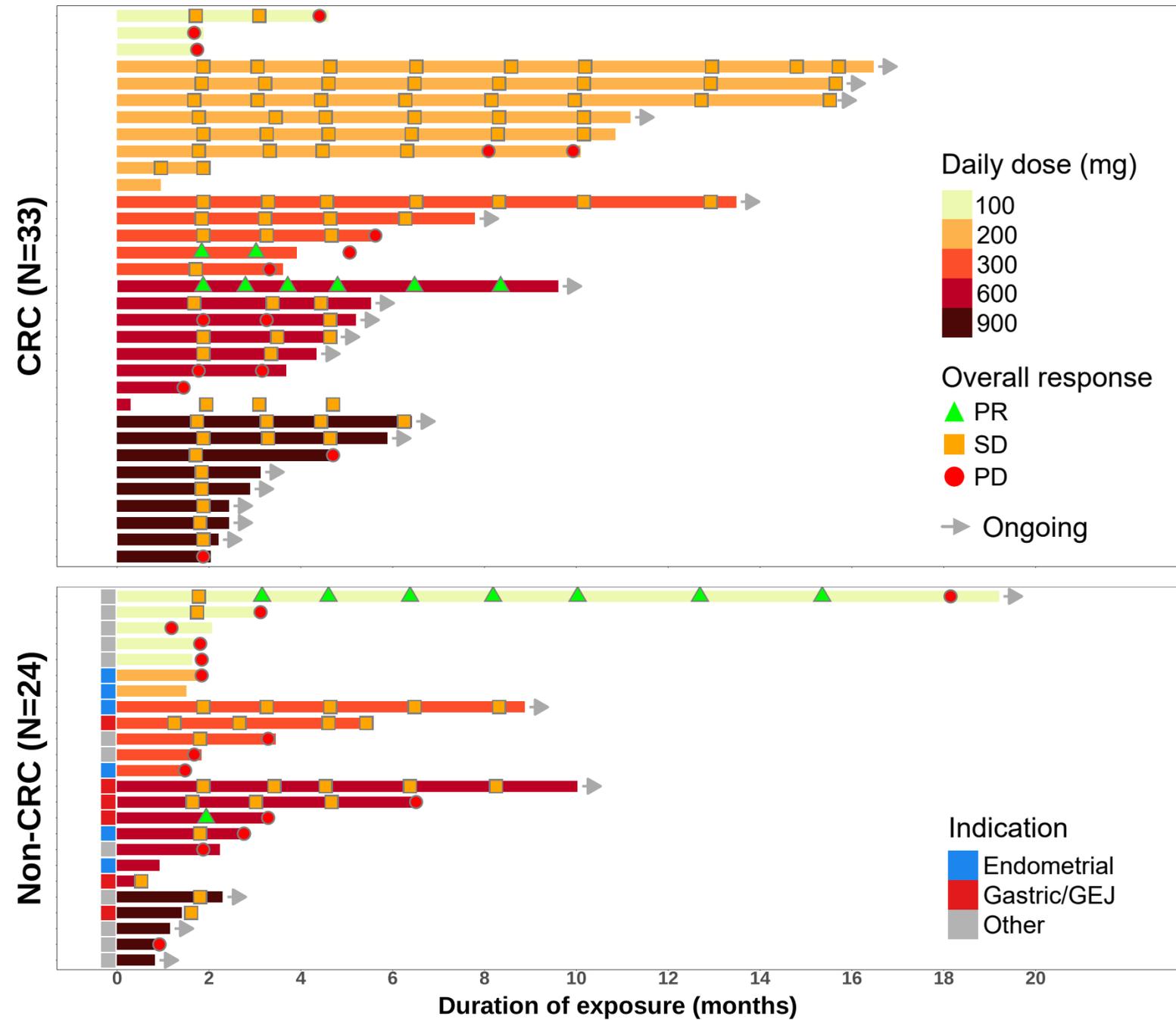
	CRC N=33	Non-CRC N=22	Total N=55*
BOR, n (%)			
PR	2 (6.1)	1 (4.5)	3 (5.5)
SD	24 (72.7)	10 (45.5)	34 (61.8)
PD	6 (18.2)	8 (36.4)	14 (25.5)
NE	1 (3.0)	3 (13.6)	4 (7.3) <sup>†</sup>
ORR, n (%)	2 (6.1)	1 (4.5)	3 (5.5)
(95% CI)	(0.7, 20.2)	(0.1, 22.8)	(1.1, 15.1)
DCR <sup>‡</sup> , n (%)	26 (78.8)	11 (50.0)	37 (67.3)
(95% CI)	(61.1, 91.0)	(28.2, 71.8)	(53.3, 79.3)

\*2 ongoing patients were excluded as they had not yet reached the first post-baseline assessment at data cutoff.  
<sup>†</sup>The 4 NE patients are not shown in the plot.  
<sup>‡</sup>DCR = PR+SD. BOR of SD is defined as at least one SD observed 6 weeks or later from the first dose, without having a better response.

BOR, best overall response with confirmation; CI, confidence interval; CRC, colorectal cancer; DCR, disease control rate; GEJ, gastroesophageal junction; NE, not evaluable; ORR, overall response rate; PD, progressive disease; PR, partial response; SD, stable disease.

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# Prolonged treatment duration and PFS in CRC patients



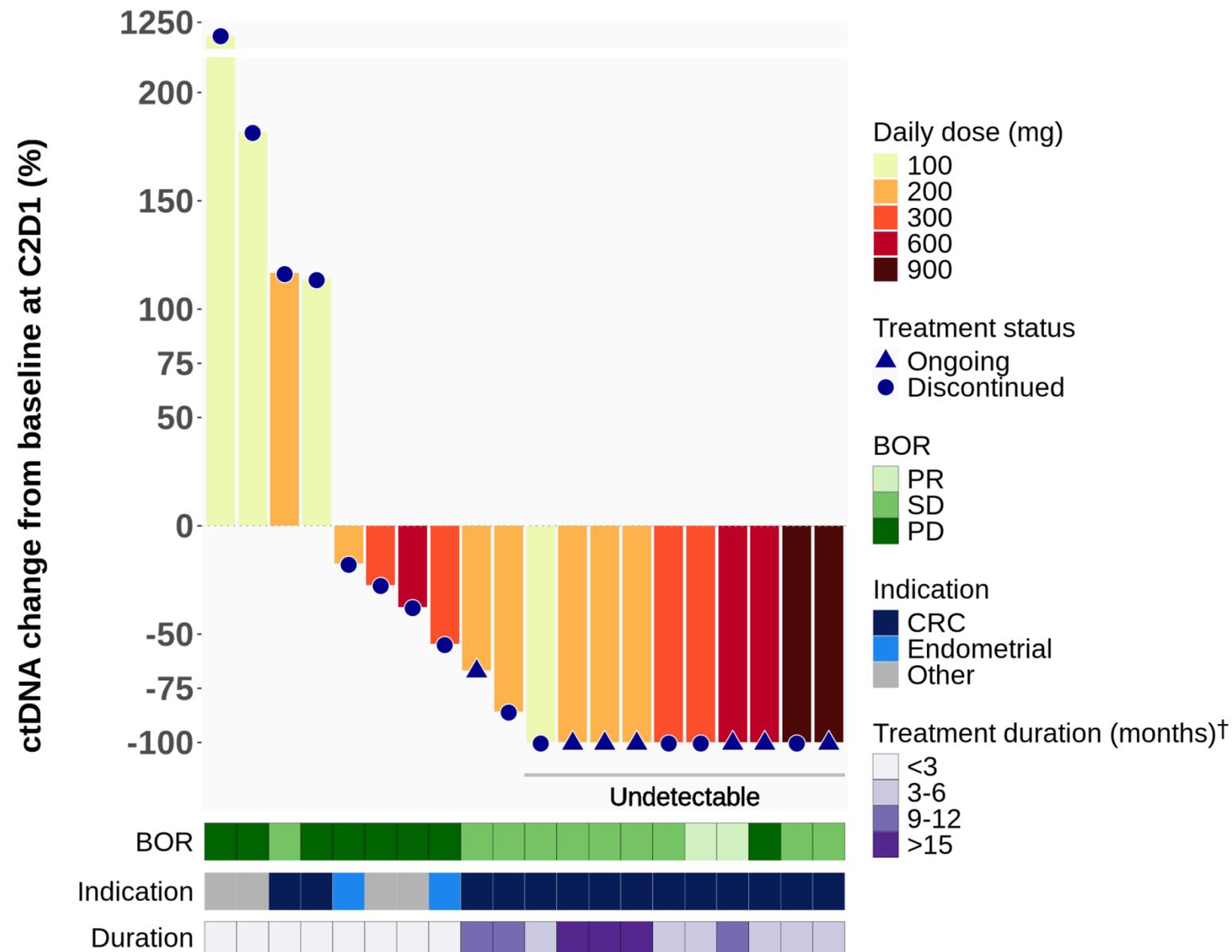
**Patients at risk**

	0	2	4	6	8	10	12	14	16	18	20
<b>CRC</b>	33	20	18	10	8	6	4	3	0	0	0
<b>Non-CRC</b>	22	12	5	4	3	1	1	1	1	1	0

\*Median PFS was not reached for the 30 CRC patients (18 ongoing) who received daily doses  $\geq 200$  mg.

CI, confidence interval; CRC, colorectal cancer; GEJ, gastroesophageal junction; mo, months; NR, not reached; PD, progressive disease; PFS, progression-free survival; PR, partial response; SD, stable disease.

# ~70% of CRC patients with detectable ctDNA at baseline cleared after ~1 month on HRO761



20 patients analyzed\*  
 ↳ 10 cleared ctDNA (all CRC, 2 PR, 7 SD, 1 PD)  
 ↳ 6 ongoing  
 ↳ 3 on treatment for > 15 months

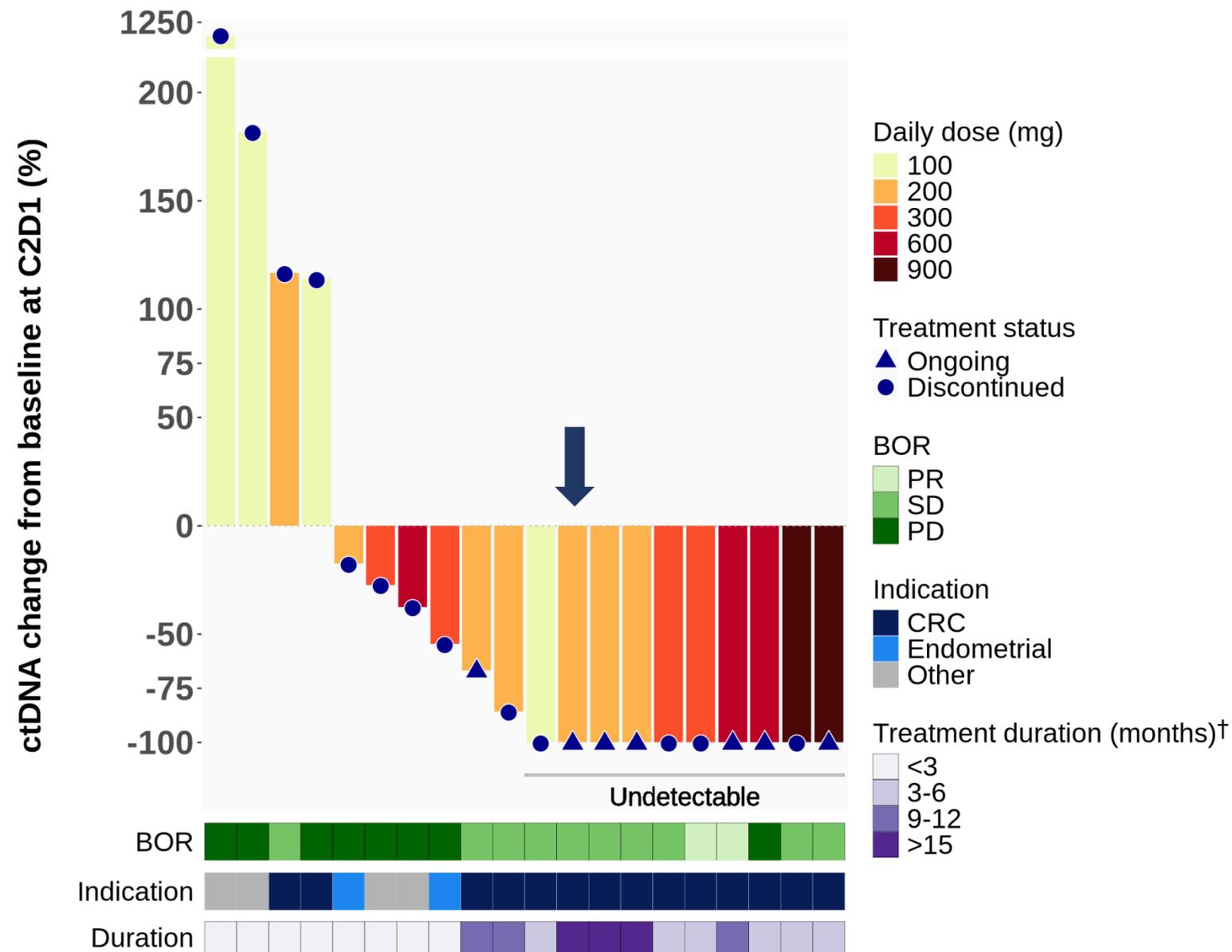
\*Analysis performed for patients with detectable ctDNA at baseline for which a C2D1 measurement was available at data cutoff (n=20, no gastric/GEJ).

†There were no patients with treatment duration of 6-9 or 12-15 months at data cutoff; therefore, these categories were excluded from the plot.

BOR, best overall response with confirmation; C2D1, day 1 of cycle 2; CEA, carcinoembryonic antigen; CRC, colorectal cancer; ctDNA, circulating tumor DNA; GEJ, gastroesophageal junction; PD, progressive disease; PR, partial response; SD, stable disease.

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# ~70% of CRC patients with detectable ctDNA at baseline cleared after ~1 month on HRO761



**12.6 x 5.4 cm**

126.0 mm  
54.2 mm

A 53-year-old woman with stage IV MLH1/PMS2-deficient transverse CRC with PD on 5 lines of therapy

↓ 1.8 months HRO761

**10.3 x 4.5 cm**

102.5 mm  
44.5 mm

- CEA 46 to 4.9
- ctDNA cleared
- BOR = SD
- > 15 months on therapy

\*Analysis performed for patients with detectable ctDNA at baseline for which a C2D1 measurement was available at data cutoff (n=20, no gastric/GEJ).

†There were no patients with treatment duration of 6-9 or 12-15 months at data cutoff; therefore, these categories were excluded from the plot.

BOR, best overall response with confirmation; C2D1, day 1 of cycle 2; CEA, carcinoembryonic antigen; CRC, colorectal cancer; ctDNA, circulating tumor DNA; GEJ, gastroesophageal junction; PD, progressive disease; PR, partial response; SD, stable disease.

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# Conclusions

- In heavily treated post-CPI patients with advanced MSI-H/dMMR cancers, HRO761 shows:
  - A favorable safety profile
  - Dose-linear exposure with no differences between QD and BID schedules
  - Durable antitumor activity in CRC patients
- ctDNA analysis highlights deep molecular responses in CRC patients, including patients with SD by RECIST
- Data continue to mature, with over half of CRC patients on treatment at data cutoff

## Future directions

- Randomized dose optimization of single agent HRO761 at 300 mg QD and 600 mg QD is ongoing
- Investigation of HRO761 in combination with pembrolizumab or irinotecan is ongoing

BID, twice daily; CPI, checkpoint inhibitor; CRC, colorectal cancer; ctDNA, circulating tumor DNA; dMMR, mismatch repair deficiency; MSI-H, microsatellite instability-high; QD, once daily; RECIST, Response Evaluation Criteria in Solid Tumors.

## Acknowledgments

The authors would like to thank all the patients who participated in the study and their caregivers, as well as personnel at all study sites.



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